## SEQUENCE LISTING

<110> Yunn-Bor Yang Chia-Li Wei Jyh-Shing Hsu Ying-Chieh Tsai													
Mutated Penicillin Expandase and Process for Preparing 7-ADCA Using the Same													
<130> 6653-017-999													
<140> To Be Assigned <141> 2002-03-26													
<160> 1													
<170> FastSEQ for Windows Version 4.0													
<210> 1 <211> 1230 <212> DNA <213> Streptomyces clavuligerus													
<220> <221> CDS <222> (232)(1164)													
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acg acg gtg ccc acc ttc agc ctg gcc gaa ctc cag cag ggc ctg cac  Thr Thr Val Pro Thr Phe Ser Leu Ala Glu Leu Gln Gln Gly Leu His  5 10 15													
cag gac gag ttc cgc agg tgt ctg agg gac aag ggc ctc ttc tat ctg Gln Asp Glu Phe Arg Arg Cys Leu Arg Asp Lys Gly Leu Phe Tyr Leu 20 25 30													
acg gac tgc ggt ctg acc gac acc gag ctg aag tcg gcc aag gac atc Thr Asp Cys Gly Leu Thr Asp Thr Glu Leu Lys Ser Ala Lys Asp Ile 35 40 45 50													
gtc atc gac ttc ttc gag cac ggc agc gag gcg gag aag cgc gcc gtc 429 Val Ile Asp Phe Phe Glu His Gly Ser Glu Ala Glu Lys Arg Ala Val 55 60 65													
acc tcg ccc gtc ccc acc atg cgc cgc ggc ttc acc ggg ctg gag tcg  Thr Ser Pro Val Pro Thr Met Arg Arg Gly Phe Thr Gly Leu Glu Ser  70  75  80													
gag agc acc gcc cag atc acc aat acc ggc agc tac tcc gac tac tcg  Glu Ser Thr Ala Gln Ile Thr Asn Thr Gly Ser Tyr Ser Asp Tyr Ser  85  90  95													

	tgc Cys 100															573
ttc Phe 115	gag Glu	cgg Arg	atc Ile	tgg Trp	acc Thr 120	cag Gln	tac Tyr	ttc Phe	gac Asp	cgc Arg 125	cag Gln	tac Tyr	acc Thr	gcc Ala	tcc Ser 130	621
	gcg Ala															669
	Gly ggg															717
tac Tyr	ttc Phe	ccg Pro 165	cag Gln	gtc Val	ccc Pro	gag Glu	cac His 170	cgc Arg	agc Ser	gcc Ala	gag Glu	gag Glu 175	cag Gln	ccc Pro	ctg Leu	765
cgg Arg	atg Met 180	gcg Ala	ccg Pro	cac His	tac Tyr	gac Asp 185	ctg Leu	tcg Ser	atg Met	gtc Val	acc Thr 190	ctc Leu	atc Ile	cag Gln	cag Gln	813
aca Thr 195	ccc Pro	tgc Cys	gcc Ala	aac Asn	ggc Gly 200	ttc Phe	gtc Val	agc Ser	ctc Leu	cag Gln 205	gcc Ala	gag Glu	gtc Val	ggc Gly	ggc Gly 210	861
	ttc Phe				Pro											909
ggc Gly	gcc Ala	atc Ile	gcg Ala 230	acc Thr	ctg Leu	gtg Val	acc Thr	ggc Gly 235	ggc Gly	cag Gln	gtc Val	aag Lys	gcc Ala 240	ccc Pro	cgg Arg	957
cac His	cat His	gtc Val 245	gcg Ala	gcc Ala	ccc Pro	cgc Arg	agg Arg 250	gac Asp	cag Gln	ata Ile	gcg Ala	ggc Gly 255	agc Ser	agc Ser	cgc Arg	1005
	tcc Ser 260															1053
	ccg Pro															1101
acc Thr	gcc Ala	acg Thr	ttc Phe	cag Gln 295	gat Asp	tgg Trp	atc Ile	Gly 999	ggc Gly 300	aac Asn	tac Tyr	gtg Val	aac Asn	atc Ile 305	cgc Arg	1149
	cgc aca tcc aag gca tagagagcac acaccgtcat ggtcacagca gcaatcagtg Arg Thr Ser Lys Ala 310													1204		
gta	ccga	cga ·	gata	cgcg	cg a	gggc	g									1230

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